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Val Leu Ser Gly Leu Ile Leu Thr Ala Thr Val Ile Ser Leu Ile Ile
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Ala Thr Pro Leu Leu Val Ile Phe Ser Pro Val Leu Val Pro Ala Leu 65 70 75 80

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Ser Lys Val Leu Ala Val Met Thr Ala Leu Pro Ile Gly Gly Thr Leu 50 55 60

Leu Ala Leu Ala Gly Ile Thr Leu Ala Gly Thr Met Ile Gly Leu Ala 65 70 75 80

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Val Cys Glu Gln Gln Ile Gln Gln Arg Asp Phe Leu Arg Ser Cys Gln 50 55 60

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Phe Val Thr Thr Leu Ser Asn Cys Ala Pro Arg Cys Thr Met Pro Gly
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Asp Leu Lys Arg Ala Ile Gly Gln Met Arg Gln Glu Ile Gln Gln Gln 115 120 125

Gly Gln Gln Gln Gln Gln Gln Glu Val Gln Arg Trp Ile Gln Gln 130 135 140

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Gln Ala Glu Ala Gly Thr Ile Glu Val Trp Asp Gln Asn Arg Gln Gln 50 55 60

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Leu Gln Ser Glu Asn Asp Asn Arg Gly Gln Ile Val Arg Val Glu Gly 65 70 75 80

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Glu Gln Gly Gly Arg Gly Gly Gly Arg Tyr Tyr Ser Asn Gly Val Glu 100 105 110

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taggtacctg gcaggtaaag actctgctc
                                                                    29
```

```
<210> 25
<211> 6
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic KpnI site
<400> 25
ggtacc
```